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Phe Lys Pro Ser Glu Thr Val Phe Lys Ile Val Phe Trp Leu Gly Tyr																				
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cta aac agc tgc atc aac ccc atc ata tac cca tgc tcc agc caa gag																				569
Leu Asn Ser Cys Ile Asn Pro Ile Ile Tyr Pro Cys Ser Ser Gln Glu																				
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Phe Lys Lys Ala Phe Gln Asn Val Leu Arg Ile Gln Cys Leu Arg Arg																				
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Gln Ala Val Glu Gly Gln His Lys Asp Met Val Arg Ile Pro Val Gly	
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Ser Arg Glu Thr Phe Tyr Arg Ile Ser Lys Thr Asp Gly Val Cys Glu	
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Trp Lys Phe Phe Ser Ser Met Pro Arg Gly Ser Ala Arg Ile Thr Val	
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tcc aaa gac caa tcc tcc tgt acc aca gcc cgg gtg aga agt aaa agc	857
Ser Lys Asp Gln Ser Ser Cys Thr Thr Ala Arg Val Arg Ser Lys Ser	
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Phe Leu Gln Val Cys Cys Cys Val Gly Pro Ser Thr Pro Ser Leu Asp	
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Lys Asn His Gln Val Pro Thr Ile Lys Val His Thr Ile Ser Leu Ser	
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Glu Asn Gly Glu Glu Val	
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 Pro Cys Ser Ser Gln Glu Phe Lys Lys Ala Phe Gln Asn Val Leu Arg
 35 40 45
 Ile Gln Cys Leu Arg Arg Lys Gln Ser Ser Lys His Ala Leu Gly Tyr
 50 55 60
 Thr Leu His Pro Pro Ser Gln Ala Val Glu Gly Gln His Lys Asp Met
 65 70 75 80
 Val Arg Ile Pro Val Gly Ser Arg Glu Thr Phe Tyr Arg Ile Ser Lys
 85 90 95
 Thr Asp Gly Val Cys Glu Trp Lys Phe Phe Ser Ser Met Pro Arg Gly
 100 105 110
 Ser Ala Arg Ile Thr Val Ser Lys Asp Gln Ser Ser Cys Thr Thr Ala
 115 120 125
 Arg Val Arg Ser Lys Ser Phe Leu Gln Val Cys Cys Cys Val Gly Pro
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 His Thr Ile Ser Leu Ser Glu Asn Gly Glu Glu Val
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 cttgtacccc tattttaaca ganaaaaaat ctgatcattg tanggagntt aaggacttgc 180
 ccaaggccaa ggggacccat gatttaaacc tgggtctcct attctcaacn tgcacatttt 240
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 ggttcctgcc cagtcgagtc tagacattnc tggggcaccc tttaagtggc ctccagcacc 240
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 ggctgacggg gtagtcagcc atgctcggag gctgaatttg gcagggactt gctgccatct 360
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 Gly His
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 Thr Pro Met Thr
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